



SEQUENCE LISTING

<110> Wilkinson, Jack
McBride, Kevin
Bertain, Sean

<120> GENETIC CONSTRUCTS HAVING HETEROLOGOUS 3' POLYADENYLATION SIGNAL SEQUENCE
MOTIFS THAT FUNCTION IN PLANTS

<130> 0325.210

<140> US 10/600,230

<141> 2003-06-20

<150> 60/390,529

<151> 2002-06-20

<160> 81

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<213> *Saccharomyces cerevisiae*

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aacacgcgtt ctgtctcttt ccaagggact ccgaatatgc cactatttat ctgtggcatt 300
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<211> 541

<212> DNA

<213> *Saccharomyces cerevisiae*

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cgaagaaaag aggaaaacgc aagtggataa aggggtgggg ggcaaaagta ttaagaaaa 480
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<211> 666
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<213> *Saccharomyces cerevisiae*

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<220>
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<400> 6
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<211> 37
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<223> PCR primer

<400> 7

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<210> 8

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 8

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<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 9

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<210> 10

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 10

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<210> 11

<211> 35

<212> DNA

<213> Artificial Sequence

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<210> 12

<211> 43

<212> DNA

<213> Artificial Sequence

<220>

<223> Mutagenic Oligonucleotide

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<210> 13
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> PCR primer

<400> 13
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 <211> 44
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<220>
 <223> Mutagenic oligonucleotide

<400> 14
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<210> 15
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Mutagenic Oligonucleotide

<400> 15
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<210> 16
 <211> 510
 <212> DNA
 <213> *Saccharomyces cerevisiae*

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 aaaaaataaa agatagaaaa gatcttagga acggatagag gtttgaaaaa ggaataacag 180
 gtaatttttc attttcatat cggttgtaac attataaagc tcacaaattt aaaacaaaaa 240
 aaaacataaa cctaacaagg ttaatcattt gcacatgac tcatcatata gatcaattca 300
 taatctatat aataatgaat aattagaata aaaatttcct cttgtctcag aacgcccac 360
 ggatggcata acttttagtta atgatcacac gacggacgaa gtattgaaag acaacctaac 420
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 gatatgaaca acctaactca caaaatttac 510

<210> 17
 <211> 877
 <212> DNA
 <213> *Saccharomyces cerevisiae*

<400> 17

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ctgtgtttgg ttttagcaca ctttccaata accaagttgg tttcagatca tccccatatt 660
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gtacagtggg ccattttttt ctgattcttc atattttccg ttataagtct tataaggaag 780
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<210> 18

<211> 669

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 18

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ccttctatat attatttctc taacagctat gttaacatga ttgcctttgt ttatctacta 180
aaggaccctt ctactttatc taccatacgc ctatatcttc tctgtgtttc aatcatatcg 240
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tcctcagagg ttttgtcaag tgggtgttgt gtgcaaactc gaagagaata gttattttatt 480
ttggcaggcg cacttgaggt tgaaagttgt agattatgtg gggatacaaa gccatttgtc 540
gagtttcgat cttccattga taacttttgt atcgacgaat atgaatcgtt aaaacgttcc 600
gtctttgtct gagaagattt ttggcctttg agagttcttt tttccctggt ataatacaaa 660
tcttcactt 669
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<210> 19

<211> 443

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 19

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ggatacagtg agcctgaaga ggacaagctg cttccatgtt gtagtgttta gatatatgag 300
cttaaaatct agatttactg aatattatac aatagtaatt atacataaag aaattccatt 360
ttatctgttc gatagcaatg gaagaggaga gagttctgtg aaacaaataa cagcagcaca 420
gaaaactccc gtcaacgtaa tat 443
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<210> 20

<211> 427

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 20

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atgttggttt	tcaagagctt	ggatttttga	tcgtcttata	ctatgacgtt	cactattttc	240
gcgaacccgg	gtaataccat	tagctatttt	gatagaaagg	gattttttatt	aggggaatata	300
accacattta	aagtgtccta	tcatgtttca	atctccagta	aacgcacata	agccgaccaa	360
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gggaaaa						427

<210> 21

<211> 810

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 21

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tctctgacac	gactcatatt	ggccctcctc	agcatcatca	tgatccattt	gggaacacct	720
tgttttacag	taatcaaccg	ttcagtaact	aagaccttac	cttgatcctt	caattctctt	780
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<210> 22

<211> 763

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 22

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tatttccgca	gctccgtagt	taataaactg	ttttaatatg	acctcaagg	tattcatata	540
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atttatacca	cagccaagta	gcaaacaata	tttattgttt	atgaagtggg	tattaactaa	720
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<210> 23

<211> 498

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 23

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tattgtttat	tttggtaaaa	tatagacgca	acttccttat	tataaagaaa	ggcattat	180
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atataaagta	actgccac					498

<210> 24

<211> 492

<212> DNA

<213> *Pichia pastoris*

<400> 24

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cagagtacag	aagattaagt	gagaccttcg	tttggtcgga	ttccccacac	accatagctt	360
caaatgttt	ctactccttt	tttactcttc	cagattttct	cggactccgc	gcacgcccgt	420
accacttcaa	aacacccaag	cacagcatac	taaattttcc	ctctttcttc	ctctaggggtg	480
tcgttaatta	cc					492

<210> 25

<211> 876

<212> DNA

<213> *Pichia pastoris*

<400> 25

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aagatacgca	tgcatataca	tatatacact	agctaacatc	cacccaatat	atataccct	720
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<210> 26

<211> 577

<212> DNA

<213> *Pichia pastoris*

<400> 26

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gcctgtgtct	cggtgtccaga	ccccgcgcgt	ccttggtgtg	agtctctgta	cgtatgggtt	180
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ggaatacaga	accctgtaga	gttaaggagt	gtaaacaccc	gacacagtat	ataccaggcc	420
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aataatgata	accagcagcg	catagcccag	tacgaggcct	tgacgtcaag	gtcagtttct	540
gcagaacaat	cgcattatcg	aatccatgga	atgcact			577

<210> 27

<211> 650

<212> DNA

<213> *Pichia pastoris*

<400> 27

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tgaccttgca	acattgtctc	ccagcgcgtt	gccaaagcga	acttgatata	agtatagtat	600
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<210> 28

<211> 412

<212> DNA

<213> *Homo sapiens*

<400> 28

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acaacactcc	aataaacatt	ttgattttag	gttctgcctc	tgagtttatt	cctgagggga	240
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aaggaaaaag	gtcactggct	agtgtagcta	gtgtaaacag	gaccagggcg	atgcatggga	360
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<210> 29

<211> 308

<212> DNA

<213> *Homo sapiens*

<400> 29

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cggagctggg	ctgcagctgg	ggctggcatg	gactttcatt	tcagagattc	ggtttttaag	180
aagatgcatg	cctagcgtgt	tctttttttt	ttccaatgat	ttgtaataata	cattttatga	240
ctggaaactt	ttttgtacaa	cactccaata	aacattttga	tttttaggttc	tgccctctgag	300
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<210> 30

<211> 363

<212> DNA

<213> *Homo sapiens*

<400> 30
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<210> 31
 <211> 341
 <212> DNA
 <213> Homo sapiens

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<211> 3107

<212> DNA

<213> Homo sapiens

<400> 68

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<210> 69

<211> 2878

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 69

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<211> 3379

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 70

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<211> 3233

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 71

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<210> 72

<211> 1775

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 72

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<210> 73

<211> 967

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 73

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<210> 74

<211> 2010

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 74

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<210> 75

<211> 6224

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 75

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<211> 2367

<212> DNA

<213> *Aspergillus nidulans*

<400> 79

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<212> DNA

<213> *Aspergillus nidulans*

<400> 80

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<400> 81

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